BGI-141CP - 25 -

	val	_			-		_		_	_	_			_	_	70
	ı gcg ı Ala		_	_	_			_	~ 5	_				_	-	144
	ctc Leu 50															192
	cat His			_	_				_	_					_	240
	gat Asp	_							-		_	_	_	_	_	288
	aat Asn	-		_		_	_	_	-		_	_	-	•	-	336
	gag Glu			_				_			_	_	_	_		384
	ggc Gly 130			_	_	_				_		_			_	432
	gta Val	_					_	_			_	_		-	_	480
	a aaa 6 Lys			_	_	_			_	_	_	_		-	_	528
	g ctt Leu				_				_			_			_	576
	g acg I Thr		-		_	_		_			_					624
_	gga Gly 210		_		_			_						_		672
_	aca J Thr				-	_			~	_		_	_			720
_	aca Thr	_		•			_	_	~	•	-		_	_		768
cct	gaa	caa	aag	cat	tcc	ttt	caa	atg	aac	cag	aca	gtg	ctt	gac	ggc	816

att gtc atg ctg acc gct tat gat tat ccg gca gct aaa ctt gct gaa 96

Pro Glu Gln Lys His Ser Phe Gln Met Asn Gln Thr Val Leu Asp Gly 260 265 270

ttg tac ggg gga aaa Leu Tyr Gly Gly Lys 275 831

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<211> 277

<212> PRT

<213> Bacillus subtilis

<400> 24

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Ile Val Met Leu Thr Ala Tyr Asp Tyr Pro Ala Ala Lys Leu Ala Glu 20 25 30

Gln Ala Gly Val Asp Met Ile Leu Val Gly Asp Ser Leu Gly Met Val
35 40 45

Val Leu Gly Leu Asp Ser Thr Val Gly Val Thr Val Ala Asp Met Ile 50 55 60

His His Thr Lys Ala Val Lys Arg Gly Ala Pro Asn Thr Phe Ile Val 65 70 75 80

Thr Asp Met Pro Phe Met Ser Tyr His Leu Ser Lys Glu Asp Thr Leu 85 90 95

Lys Asn Ala Ala Ile Val Gln Glu Ser Gly Ala Asp Ala Leu Lys
100 105 110

Leu Glu Gly Glu Gly Val Phe Glu Ser Ile Arg Ala Leu Thr Leu 115 120 125

Gly Gly Ile Pro Val Val Ser His Leu Gly Leu Thr Pro Gln Ser Val 130 135 140

Gly Val Leu Gly Gly Tyr Lys Val Gln Gly Lys Asp Glu Gln Ser Ala 145 150 150 160

Lys Lys Leu Ile Glu Asp Ser Ile Lys Cys Glu Glu Ala Gly Ala Met 165 170 175

Met Leu Val Leu Glu Cys Val Pro Ala Glu Leu Thr Ala Lys Ile Ala 180 185 190

Glu Thr Leu Ser Ile Pro Val Ile Gly Ile Gly Ala Gly Val Lys Ala 195 200 205

Asp Gly Gln Val Leu Val Tyr His Asp Ile Ile Gly His Gly Val Glu 210 220

Arg Thr Pro Lys Phe Val Lys Gln Tyr Thr Arg Ile Asp Glu Thr Ile 225 230 235 240

Glu Thr Ala Ile Ser Gly Tyr Val Gln Asp Val Arg His Arg Ala Phe 245 250 255 260

165

270

Pro Glu Gln Lys His Ser Phe Gln Met Asn Gln Thr Val Leu Asp Gly

265

			200										2.0			
Leu	Tyr	Gly 275	Gly	Lys												
<211 <212	0> 25 L> 85 2> DN 3> Ba	8 1A	lus :	subti	ilis											
)> L> CI 2> (1		(858))												
atg	0> 25 aga Arg	cag														48
	cat His									•	_	_	_			96
	cat His															144
_	gcc Ala 50	_		_	-						_					192
	gaa Glu	-		_	_		_	_	-				_	_	_	240
	gca Ala	_		_		_	_				_		_	_		288
-	atg Met				_	_		_	_			_	_	~	_	336
	gac Asp			_		_		_	-	• -			-		_	384
	atc Ile 130	_	-	_	_					_	_	_		-	gcc Ala	432
	ttc Phe				_		_		-	•	-	-	-		tta Leu 160	480
	agc Ser								-	-		_	_	_	_	528

170

175